My suggestions for session chairs.

scott

Systems Biology & Applied Genomics I—Jesse Poland	Session Chair?		
Genomic Assisted Breeding & Integrated Platforms I—Rajeev VArshney	Session Chair?		
Evolution Selection—Jianming Yu	Session Chair?		
Genomic Assisted Breeding & Integrated Platforms II—Esther van der Knaap			
Functional Validation—CJ TSAI			
Systems Biology & Applied Genomics II—Gerald Tuskan			Session
Genome Editing and Targeted Modifications—Todd Mockler		Se SessionC	essionChair? hair?
Developing World Crops—Jean Marcel Ribaut	90 Min	SessionChair? SessionChair?	
Roundtable: Applying Genomic Techniques to Developing World Germplasm		Chair?	
On Apr 20, 2015, at 4:45 PM, Peggy Ozias-Akins <pre>cpozias@uga.edu></pre> wrote:			
This looks good to me.			
Peggy			
On Apr 21, 2015, at 6:25 AM, Jeremy Schmutz <jschmutz@hudsonalpha.org> wrote:</jschmutz@hudsonalpha.org>			
Deborah - how about this? 10 minutes means half time. Scott and Peggy?			

Jeremy

Evolution/Selection:

John Vogel- A Quartet of Brachypodium Species Facilitate the Discovery and Characterization of Agriculturally Relevant Grass Genes

10 minutes: Justin Vaughn The Genomic Signature of a Century of North American Soybean Improvement 10 minutes: Wolfgang Goettel Comparative Seed Transcriptome Analysis of Landraces and Milestone Cultivars Representing 90 Years of North American Soybean Breeding

Genomic Assisted Breeding & Integrated Platforms II:

10 minutes: Stephanie Verhoff Development of Nearly Isogenic Lines to Study Partial Resistance to Phytophthora Stem and Root Rot 10 minutes: Sandra Truong Crop Improvement by Genetic Variation in Leaf Angle

Functional Validation:

Pamela Pena: Modulating Nitrogen Flux in Sorghum and Wheat

Soraya C. M. Leal-bertioli : Identification of QTLs for Rust Resistance in the Wild Peanut Relative Arachis Magna and the Development Markers for Introgression of this Resistance into Cultivated Peanut

Systems Biology & Applied Genomics II:

Phil McClean - Genomic Perspectives on Common Bean Improvement

Genome Editing:

CJ Tsai - Exploiting SNPs for Efficient CRISPR Mutations in Highly Heterozygous Outcrossing Species

On Apr 20, 2015, at 3:02 PM, Deborah Powell wrote:

All,

Are we getting close on assigning the program speakers from the poster abstracts that were submitted? We are working on a tight timeline for getting the conference program to print. Once the selected posters have been assigned an oral speaking slot, we will need to contact the speaker to get their acceptance, rearrange the conference program and agenda to accommodate the abstracts of the oral speakers, edit and get approval on the conference program guide and have it ready to print on Monday, April 27.

Thanks,

Deborah

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